

Sequence Listing.ST25.txt
SEQUENCE LISTING

<110> Pfizer Ltd. (EP (GB) only)
Pfizer Inc. (US, JP, EP except GB)
Fidock, Mark David

<120> Novel Polypeptide

<130> PC10960AGPR

<150> GB 0030855.1

<151> 2000-12-18

<150> GB 0101222.8

<151> 2001-01-17

<160> 7

<170> PatentIn version 3.0

<210> 1

<211> 1082

<212> DNA

<213> Homo sapiens

<400> 1

atgctgtcca ttttgcttcc ttccagggga agcagaagcg ggagccgtcg tggagctctg	60
ctcctggagg gagcctcccg ggacatggag aaggtggaca tgaatacatc acaggaacaa	120
ggtctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc	180
atctttatcc tagggctgcc actaaatggc actgtcttgt ggcactcctg gggccaaacc	240
aagcgctgga gctgtgccac cacctatctg gtgaacctga tgggtggccga cctgctttat	300
gtgctattgc ccttcctcat catcacctac tctactagatg acaggtggcc cttcggggag	360
ctgctctgca agctggtgca cttcctgttc tatatcaacc tttacggcag catcctgctg	420
ctgacctgca tctctgtgca ccagttccta ggtgtgtgcc acccaactgtg ttcgctgccc	480
taccggaccc gcaggcatgc ctggctgggc accagcacca cctgggccct ggtggtcctc	540
cagctgctgc ccacactggc cttctccac acggactaca tcaatggcca gatgatctgg	600
tatgacatga ccagccaaga gaattttgat cggctttttg cctacggcat agttctgaca	660
ttgtctggct ttctttccct ccttggtcat tttggtgtgc tattcactga tggtcaggag	720
cctgatcaag ccagaggaga acctcatgag gacaggcaac acagcccag ccaggtccat	780
ccggaccatc ctactggtgt gtggcctctt caccctctgt tttgtgccct tccatatcac	840

Sequence Listing.ST25.txt

```
tcgctccttc tacctcacca tctgctttct gctttctcag gactgccagc tcttgatggc 900
agccagtgtg gcctacaaga tatggaggcc tctggtgagt gtgagcagct gcctcaaccc 960
agtctgttac tttctttcaa ggggggcaaa aatagagtca ggctcctcca gaaactgagg 1020
cagaacaagt tgggtgagca tccagctggg aggaagagat gcccagggtt gaacagatct 1080
gg 1082
```

```
<210> 2
<211> 360
<212> PRT
<213> Homo sapiens
```

```
<400> 2
```

```
Met Leu Ser Ile Leu Leu Pro Ser Arg Gly Ser Arg Ser Gly Ser Arg
1 5 10 15
Arg Gly Ala Leu Leu Leu Glu Gly Ala Ser Arg Asp Met Glu Lys Val
20 25 30
Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys
35 40 45
Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu
50 55 60
Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr
65 70 75 80
Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala
85 90 95
Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu
100 105 110
Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe
115 120 125
Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile
130 135 140
Ser Val His Gln Phe Leu Gly Val Cys His Pro Leu Cys Ser Leu Pro
145 150 155 160
Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala
165 170 175
Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp
```

Sequence Listing.ST25.txt

180

185

190

Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn
195 200 205

Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe
210 215 220

Leu Ser Leu Leu Gly His Phe Gly Val Leu Phe Thr Asp Gly Gln Glu
225 230 235 240

Pro Asp Gln Ala Arg Gly Glu Pro His Glu Asp Arg Gln His Ser Pro
245 250 255

Ser Gln Val His Pro Asp His Pro Thr Gly Val Trp Pro Leu His Pro
260 265 270

Leu Phe Cys Ala Leu Pro Tyr His Ser Leu Leu Leu Pro His His Leu
275 280 285

Leu Ser Ala Phe Ser Gly Leu Pro Ala Leu Asp Gly Ser Gln Cys Gly
290 295 300

Leu Gln Asp Met Glu Ala Ser Gly Glu Cys Glu Gln Leu Pro Gln Pro
305 310 315 320

Ser Pro Val Leu Ser Phe Lys Gly Gly Lys Asn Arg Val Arg Leu Leu
325 330 335

Gln Lys Leu Arg Gln Asn Lys Leu Gly Glu His Pro Ala Gly Arg Lys
340 345 350

Arg Cys Pro Gly Leu Asn Arg Ser
355 360

<210> 3

<211> 1020

<212> DNA

<213> Homo sapiens

<400> 3

atgctgtcca ttttgcttcc ttccagggga agcagaagcg ggagccgtcg tggagctctg 60

ctcctggagg gagcctcccg ggacatggag aaggtggaca tgaatacatc acaggaacaa 120

ggctctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc 180

atctttatcc tagggctgcc actaaatggc actgtcttgt ggcaactcctg gggccaaacc 240

aagcgctgga gctgtgccac cacctatctg gtgaacctga tgggtggccga cctgctttat 300

gtgctattgc ccttcctcat catcacctac tcactagatg acaggtggcc cttcggggag 360

Sequence Listing.ST25.txt

```

ctgctctgca agctggtgca cttcctgttc tatatcaacc tttagcgagcag catcctgctg 420
ctgacctgca tctctgtgca ccagttccta ggtgtgtggc acccactgtg ttcgctgccc 480
taccggaccc gcaggcatgc ctggctgggc accagcacca cctgggcccct ggtggtcctc 540
cagctgctgc ccacactggc cttctccac acggactaca tcaatggcca gatgatctgg 600
tatgacatga ccagccaaga gaattttgat cggctttttg cctacggcat agttctgaca 660
ttgtctggct ttctttcccc ctcttggtc attttggtgt gctattcact gatggtcagg 720
agcctgatca agccagagga gaacctcatg aggacaggca acacagcccg agccaggctc 780
atccggacca tctactggt gtgtggcctc ttcaccctct gttttgtgcc ctcccatatc 840
actcgctcct tctacctcac catctgcttt ctgctttctc aggactgcca gctcttgatg 900
gcacccagtg tggcctacaa gatatggagg cctctggtga gtgtgagcag ctgcctcaac 960
ccagtcctgt actttctttc aaggggggca aaaatagagt caggctcctc cagaaactga 1020

```

```

>210> 4
>211> 339
>212> PRT
>213> Homo sapiens
>400> 4

```

```

Met Leu Ser Ile Leu Leu Pro Ser Arg Gly Ser Arg Ser Gly Ser Arg
1 5 10 15
Arg Gly Ala Leu Leu Leu Glu Gly Ala Ser Arg Asp Met Glu Lys Val
20 25 30
Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys
35 40 45
Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu
50 55 60
Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr
65 70 75 80
Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala
85 90 95
Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu
100 105 110
Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe
115 120 125

```

Sequence Listing.ST25.txt

Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile
 130 135 140
 Ser Val His Gln Phe Leu Gly Val Trp His Pro Leu Cys Ser Leu Pro
 145 150 155 160
 Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala
 165 170 175
 Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp
 180 185 190
 Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn
 195 200 205
 Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe
 210 215 220
 Leu Ser Pro Ser Leu Val Ile Leu Val Cys Tyr Ser Leu Met Val Arg
 225 230 235 240
 Ser Leu Ile Lys Pro Glu Glu Asn Leu Met Arg Thr Gly Asn Thr Ala
 245 250 255
 Arg Ala Arg Ser Ile Arg Thr Ile Leu Leu Val Cys Gly Leu Phe Thr
 260 265 270
 Leu Cys Phe Val Pro Phe His Ile Thr Arg Ser Phe Tyr Leu Thr Ile
 275 280 285
 Cys Phe Leu Leu Ser Gln Asp Cys Gln Leu Leu Met Ala Pro Ser Val
 290 295 300
 Ala Tyr Lys Ile Trp Arg Pro Leu Val Ser Val Ser Ser Cys Leu Asn
 305 310 315 320
 Pro Val Leu Tyr Phe Leu Ser Arg Gly Ala Lys Ile Glu Ser Gly Ser
 325 330 335
 Ser Arg Asn

<210> 5
 <211> 27
 <212> DNA
 <213> Homo sapiens

<400> 5
 accatgctgt ccattttgct tccttcc

27

<210> 6

Sequence Listing.ST25.txt

<211> 24
<212> DNA
<213> Homo sapiens

<400> 6
tcaccagatc tggtcaaccc tggg

24

<210> 7
<211> 24
<212> DNA
<213> Homo sapiens

<400> 7
tcagtttctg gaggagcctg actc

24